Amendments to the Specification:

The paragraph numbers of Patent Application Publication No. 2002/0009736, the published version of the present application as originally filed, will be used herein to describe the locations of the requested amendments to the specification.

Please replace paragraph [0025] of the present application with the following paragraph:

There are several DNA microchip technology reviews in the literature (Bowtell, D.D.L. Nature Genetics Supplement 21:25-32 (1999); Constantine and Herrington, Life Science News 1:11-13 (1998); Ramsay, G. Nature Biotechnology 16:40-44 (1998)), and several good web sties detailing the apparatus and protocols used by other laboratories. Table 1 lists several good web sites for organizations and entities, including highly active laboratories in DNA microchip technology, as well as several sources of robotics systems and equipment imaging software and systems and vendors of robotic components, each of which have an associated web site containing useful information.

Please replace Table 1 of page 4 of the present application with the following with the following Table:

Table 1: Informative web sites for DNA microarray technology

DNA microarray technology web sites organizations and entities

Automation and Miniaturization in Genome Analysis,

Max Plank Institute for Molecular Genetics

http://www.mpimg-berlin-dahlem.mpg.de/- autom/autom.htm

Department of Molecular Biotechnology,

University of Washington

http://chroma.mbt.washington.edu/mod - www/

Functional Genomics Group,

Albert Einstein College of Medicine

http://sequence.accom.yu.edu/bioinf/funcgenomic.html

Genomics Group,

Children's Hospital of Philadelphia

http://w95vc1.neuro.chop.edulvcheunne

Laboratory of Cancer Genetics,

National Human Genome Research Institute

http://www.nhgri.nih.govllntramural JosearchlLab cancer/

Joint Genome Institute,

Lawrence Livermore National Laboratory

http://IlnJ.gov/automation-robotics/poster.l.html

Pat Brown Laboratory,

Stanford University

http://emgm. stanford. edu/pbrown

Stanford DNA sequence and Technology Center

Stanford University

http://-sequence.stanford.edu/group/techdev/

Microarrayers. imaging systems and scanners

Applied Scientific Instrumentation, Inc.

http://www.ASIimaging.com/

Axon Instruments, Inc.

http://axon.com/GN_Genomics.html

Beecher Instruments

http://www.beecherinstruments.com/

BioDiscovery, Inc.

http://www.Biodiscovery.com/

BioRobotics, Ltd.

http://www.biorobotics.com/

Empix Imaging, Inc.

http://www.empix.com/

GeneMachines, Genomic Instrumentation Services, Inc.

http://www.genemachines.com/

General Microarray Information

http://www.microarray.org/

General Scanning, Inc.

http://www.gensean.com/

Genetic MicroSystems, Inc.

http://www.geneticmicro.com/

Genometrix, Inc.

http://www.genometrix.com/

Genomic Solutions

http://www.genomicsolutions.com/

Imaging Research, Inc.

http://www.imagingresearch.com/

Intelligent Automation

http://www-ias.com

Molecular Dynamics, Inc.

http://www.mdyn.com/arrays/arraywhat.htm

Radius Biosciences

http://www.ultranot.com/- radius

Research Genetics

http://www.resgen.com

ScanAlyze software

http://bronzino.stanford.edu/ScanAlyze/

Telechem International, Inc.

http://www.wenet/~telechem/

Western Technology Marketing

http://www.westerntechnology.com/

Robotics Galil

http://galilme.com/

Parker-Compumotor

http://www.compumeter.com/

Parker-Daedal

http://www.daodalpositioning.com/

Please replace paragraph [0080] of the present application with the following paragraph:

Enter your e-mail address to receive the results, which will probably take overnight. When you receive your results, go to edit and select "Find". Enter part of the known core element sequence and visually search for the second part (e.g., core element=RTGACNNNGC[SEQ ID NO:1], enter TGAC and visually search for GC 3 bases away).

Please replace the chart between paragraphs [0082] and [0083] of the present application with the following chart:

3 Feature: ID DT DT DE DE DE KW	promoter AF029342 08-APR-1998 08-APR-1998 Homo sapiens growth h gene, promoter region.	(11976) standard; DNA; HUM; 2056 BP. (Rel. 55, Created) (Rel. 55, Last updated, Version 1) commone-releasing hormone receptor				
matrix name	matrix position (str)	core simil	matrix simil.	CAGNAMAA		
/tmp/bigbox taaaaGTGAccage	1094 (+)	1.000	0.940	sequence		

Please replace the chart between paragraphs [0096] and [0097] of the present application with the following chart:

tm 60 28	gc % 50 00	any 6 00	0 00	seq TCTCCAAGTCGACA- CTTTTCC [SEQ ID NO: 3]
	60 28	60 28 50 00	60 28 50 00 6 00	60 28 50 00 6 00 0 00

SEQUENCE SIZE 1617

INCLUDED REGION SIZE 1617

PRODUCT SIZE 452, PRODUCT Tm 83 0666 PAIR ANY COMPL 6 00, PAIR 3' COMPL 2 00

PRODUCT Tm - min (OLIGO Tm) 22 8601

^{*}Notice the frequency of repeats in this sequence. It is probably not a good candidate for consideration!